

SEQUENCE LISTING

DANA-FARBER CANCER INSTITUTE, INC. KOLODNER, Richard WINAND, Nena

<120> A METHOD OF DETECTION OF ALTERATIONS IN MSH5

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<141> 1999-12-22

<150> 60/051,686

<151> 1997-07-03

<150> PCT/US98/13850

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Gly	Phe 210	Lys	Lys	Phe	Val	Leu 215	Thr	His	Leu	Val	Ser 220	Ile	Asp	Gln	Asp
Thr 225	Tyr	Ser	Val	Leu	Gln 230	Ile	Phe	Lys	Ser	Glu 235	Ser	His	Pro	Ser	Val 240
Tyr	Lys	Val	Ala	Ser 245	Gly	Leu	Lys	Glu	Gly 250	Leu	Ser	Leu	Phe	Gly 255	Ile
Leu	Asn	Arg	Суs 260	Arg	Cys	Lys	Trp	Gly 265	Gln	Lys	Leu	Leu	Arg 270	Leu	Trp
Phe	Thr	Arg 275	Pro	Thr	Arg	Glu	Leu 280	Arg	Glu	Leu	Asn	Ser 285	Arg	Leu	Asp
Val	Ile 290	Gln	Phe	Phe	Leu	Met 295	Pro	Gln	Asn	Leu	Asp 300	Met	Ala	Gln	Met
Leu 305	His	Arg	Leu	Leu	Ser 310	His	Ile	Lys	Asn	Val 315	Pro	Leu	Ile	Leu	Lys 320
Arg	Met	Lys	Leu	Ser 325	His	Thr	Lys	Val	Ser 330	Asp	Trp	Gln	Val	Leu 335	Tyr

Lys Thr Val Tyr Ser Ala Leu Gly Leu Arg Asp Ala Cys Arg Ser Leu Pro Gln Ser Ile Gln Leu Phe Gln Asp Ile Ala Gln Glu Phe Ser Asp Asp Leu His His Ile Ala Ser Leu Ile Gly Lys Val Val Asp Phe Glu Glu Ser Leu Ala Glu Asn Arg Phe Thr Val Leu Pro Asn Ile Asp Pro Asp Ile Asp Ala Lys Lys Arg Arg Leu Ile Gly Leu Pro Ser Phe Leu Thr Glu Val Ala Gln Lys Glu Leu Glu Asn Leu Asp Ser Arg Ile Pro Ser Cys Ser Val Ile Tyr Ile Pro Leu Ile Gly Phe Leu Leu Ser Ile Pro Arg Leu Pro Phe Met Val Glu Ala Ser Asp Phe Glu Ile Glu Gly Leu Asp Phe Met Phe Leu Ser Glu Asp Lys Leu His Tyr Arg Ser Ala Arg Thr Lys Glu Leu Asp Thr Leu Leu Gly Asp Leu His Cys Glu Ile Arg Asp Gln Glu Thr Leu Leu Met Tyr Gln Leu Gln Cys Gln Val Leu Ala Arg Ala Ser Val Leu Thr Arg Val Leu Asp Leu Ala Ser Arg Leu Asp Val Leu Leu Ala Leu Ala Ser Ala Ala Arg Asp Tyr Gly Tyr Ser Arg Pro His Tyr Ser Pro Cys Ile His Gly Val Arg Ile Arg Asn Gly Arg His Pro Leu Met Glu Leu Cys Ala Arg Thr Phe Val Pro Asn Ser Thr Asp Cys Gly Gly Asp Gln Gly Arg Val Lys Val Ile Thr Gly Pro

Asn Ser Ser Gly Lys Ser Ile Tyr Leu Lys Gln Val Gly Leu Ile Thr Phe Met Ala Leu Val Gly Ser Phe Val Pro Ala Glu Glu Ala Glu Ile Gly Val Ile Asp Ala Ile Phe Thr Arg Ile His Ser Cys Glu Ser Ile Ser Leu Gly Leu Ser Thr Phe Met Ile Asp Leu Asn Gln Val Ala Lys Ala Val Asn Asn Ala Thr Glu His Ser Leu Val Leu Ile Asp Glu Phe Gly Lys Gly Thr Asn Ser Val Asp Gly Leu Ala Leu Leu Ala Ala Val Leu Arg His Trp Leu Ala Leu Gly Pro Ser Cys Pro His Val Phe Val Ala Thr Asn Phe Leu Ser Leu Val Gln Leu Gln Leu Pro Gln Gly Pro Leu Val Gln Tyr Leu Thr Met Glu Thr Cys Glu Asp Gly Glu Asp Leu Val Phe Phe Tyr Gln Leu Cys Gln Gly Val Ala Ser Ala Ser His Ala Ser His Thr Ala Ala Gln Ala Gly Leu Pro Asp Pro Leu Ile Ala Arg Gly Lys Glu Val Ser Asp Leu Ile Arg Ser Gly Lys Pro Ile Lys Ala Thr Asn Glu Leu Leu Arg Arg Asn Gln Met Glu Asn Cys Gln Ala Leu Val Asp Lys Phe Leu Lys Leu Asp Leu Glu Asp Pro Thr Leu Asp Leu Asp Ile Phe Ile Ser Gln Glu Val Leu Pro Ala Ala Pro Thr Ile

Leu

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ttgggagccg ggnn
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ataagacatg gtaaacccta cacttatgag tgattctaat agtgatttcc tttcttcctt 180
gctggacag
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450

323

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<400> 61

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cattagtgtt actagttcta ttaataccat tattttgacc aaaatcctca attccagaca 480
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atattatatg tagaataaaa agagaattag actaagagtc tgaaaatttg gttcttgctc 600
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<211> 164
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ggagaatcta agggctaatg agactttggg aagaagactg ggacaatatt cagagagggg 180
gacaaaggaa gtggagttgt ggaacgaact cagactgctt cctgcttttt tgttttctgt 240
cctcag
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<221> intron
<222> (412)..(413)
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ggcgggtgga tcgcctgagg tcaggagttt gagacctgac caatatggta aaaccccgtc 360
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<221> intron
<222> (355)..(356)
<223> N = A or T or G or C
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caatccaaat ttcttaccta tttgtacccc ccgccccca agcttgagca tcttcccata 180
ctttgtggct gtacagtgtg ttgcatatca gccattactt taccaattct gtgttccttc 240
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<211> 426
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gateggtgae gteagegtee gagggaagae ggetgeeace ggeggggeea gttgagggaa 180
ctaggtagtt aagtgttgtc gggctaaaag tccctagagt gtccatccct cccccatctc 240
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catgtgcggt aatcccagct catttagggg ccaggcacca actttggttg cctttgtgcc 300

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<212> DNA
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<220>
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<222> (359)..(360)
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<210> 69
<211> 447
<212> DNA
<213> Human
<400> 69
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<210> 70
<211> 127
<212> DNA
<213> Human
<400> 70
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II	Matter 1	
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-+	TAT.	

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Junuarugua	- July Cicco	-auguauge	99-94	5449409194		00

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